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Schmickel  
1 of 15

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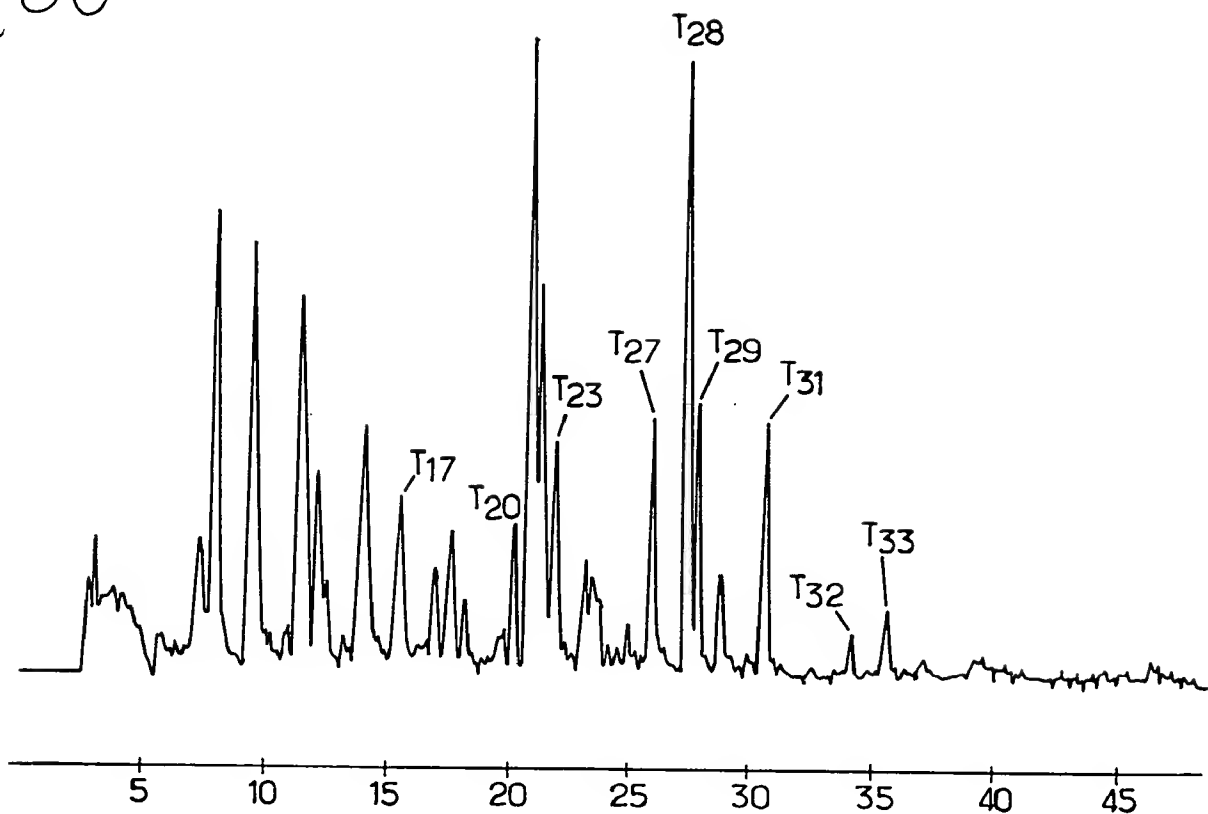


FIG. 1

Elution profile by measurement of the optical density at 218 nm  
of the product of tryptic digestion of urate oxidase

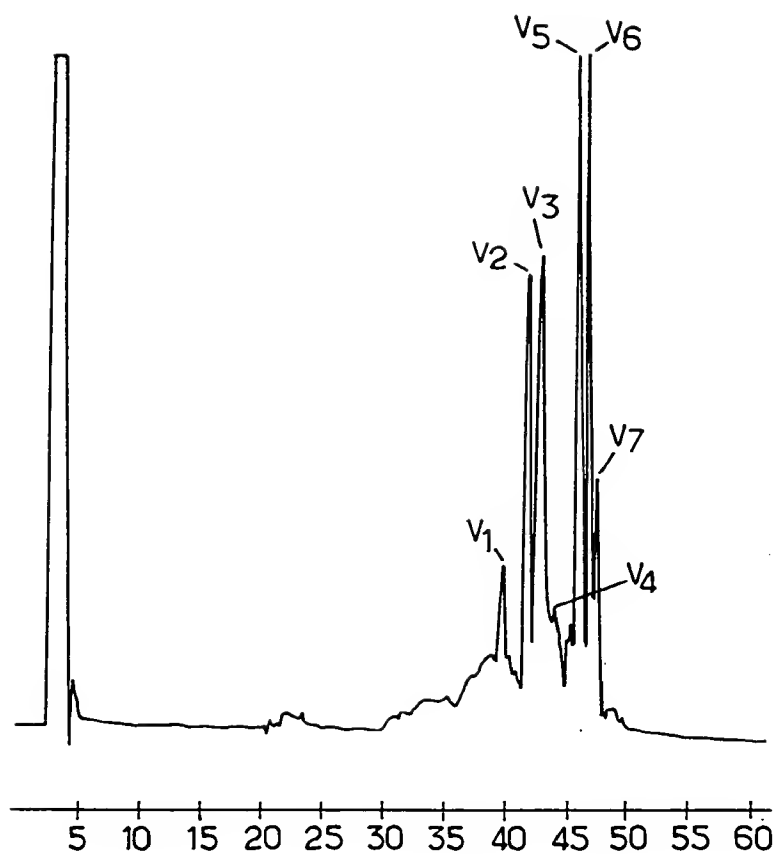


FIG. 2

Elution profile by measurement of the optical density at 218 nm  
of the product of digestion of urate oxidase with protease V8

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1  AAACCTCACTGCCTCTCTCATTCCTTCCG GTGCCCCCGATCCTCAATCCAACCTGTACA 60
61  TACTTCTCCCAACTCTCTGCTATATCCTTC ATATTCCCATACTACAAGATGTCGCCAGTA 120
121 AAAGCAGCCCGCTACGGCAAGGACAATGTC CGCGTCTACAAGGTTCAACAAGGACGAGAAG 180
181 ACCGGTGTCCAGACGGGTGACGAGATGACC GTCTGTGTGCTTCTTGAGGGTGAGATTGAG 240
241 ACCTCTTACACCAAGGGCCGACACAGCGTC ATTGTGCGAACCGACTCCATTAAAGAACACC 300
301 ATTTACATCACCGCCAAGCAGAACCCCGTT ACTCCTCCCGAGCTGTTCCGGCTCCATCCTG 360
361 GGCACACACTTCATTGAGAAGTACAACCCAC ATCCATGCCGCTCAGTCAACATTGTCTGC 420
421 CACCGCTGGACCCGGATGGACATTGACGGC AAGCCACACCCCTCACTCCTTCATCCGCGAC 480
481 AGCGAGGAGAAAGCGGAATGTGCAGGTGGAC GTGGTCGAGGGCAAGGGCATCGATATCAAG 540
541 TCGTCTCTGTCCGGCCCTGACCGTGCTGAAG AGCACCAACTCGCAGTTCTGGGGCTTCCTG 600
601 CGTGACGAGTACACCCACACTTAAGGAGACC TGGGACCGTATCCTGAGCACCCGCGTCGAT 660
661 GCCACTTGGCAGTGGAGAATTTCAAGTGA CTCCAGGAGGTCCGCTCGCACGTGCCCTAAG 720
721 TTCGATGCTACCTGGGCCACTGCTCGGAG GTCACCTCTGAAGACTTTTGTGTAAGATAAC 780
781 AGTGCCAGCGTGCAGGCCCACTATGTACAAG ATGGCAGAGCAAAATCCTGGCCGCCAGCAG 840
841 CTGATCGAGACTGTGCGAGTACTCGTTGCC T AACAAGCACTATTTTCGAAATCGACCTGAGC 900
901 TGGCACAAGGGCCCTCCAAAACACCCGGCAAG AACGCCGAGGTCTTCGCTCCTCAGTCGGAC 960
961 CCCAACGGTCTGATCAAGTGATCCGTCGGC CGGTCTCTCTGAAGTCTAAATTGTAAACC 1020
1021 AACATGATTCTCAGGTTCCGGAGTTTCCAA GGCAAACTGTATATAGTCTGGGATAGGTA 1080
1081 TAGCATTCACTTGTTTTTTTACTTCCA AAAAAAAAAA...

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# FIG. 3

Nucleotide sequence of clone 9C and of part of clone 9A

↓ : start of clone 9A

109	ATGTCCGAGTAAAGCAGCCCGCTACGGC	AAGACAATGTCCGGCTCTACAAGTTTAC	168
1	MetSerAlaValLysAlaAlaArgTyrGly	LysAspAsnValArgValTyrLysValHis	20
169	AAGGACGAGAGACCGGTGTCTCAGACGGTG	TACGAGATGACCGTCTGTGTCTCTCTGGAG	228
21	LysAspGluLysThrGlyValGlnThrVal	TyrGluMetThrValCysValLeuLeuGlu	40
229	GGTGAGATTGAGACCTCTTACACCAAGGCC	GACAACAGCGTCATTGTGCGAACCGACTCC	288
41	GlyGluIleGluThrSerTyrThrLysAla	AspAsnSerValIleValAlaThrAspSer	60
289	ATTAAGAACACCATTTACATCACCGCCAAG	CAGAACCCCGTTACTCTCCCGAGCTGTTC	348
61	IleLysAsnThrIleTyrIleThrAlaLys	GlnAsnProValThrProGluLeuPhe	80
349	GGCTCCATCCTGGGCACACACTTCATTGAG	AAGTACAACCATCTCCATGCCGCTCACGTC	408
81	GlySerIleLeuGlyThrHisPheIleGlu	LysTyrAsnHisIleHisAlaAlaHisVal	100
409	AACATTGTCTGCCACCGCTGGACCCGGATG	GACATTGACGGCAAGCCACACCCCTCACTCC	468
101	AsnIleValCysHisArgTyrThrArgMet	AspIleAspGlyLysProHisProHisSer	120
469	TTCATCCGCGACAGGAGGAGAAGCGGAAT	GTGCAGGTGGACGTGGTCGAGGGCAAGGGC	528
121	PheIleArgAspSerGluGluLysArgAsn	ValGlnValAspValValGluGlyLysGly	140
529	ATCGATATCAAGTCGTCTCTGTCCGGCCCTG	ACCGTGTCTGAAGAGACCAACTCGCAGTTC	588
141	IleAspIleLysSerSerLeuSerGlyLeu	ThrValLeuLysSerThrAsnSerGlnPhe	160
589	TGGGGCTTCCTGCGTGACGAGTACACCACA	CTTAAGGAGACCTGGGACCGGTATCCTGAGC	648
161	TrpGlyPheLeuArgAspGluTyrThrThr	LeuLysGluThrTrpAspArgIleLeuSer	180
649	ACCGACGTCGATGCCACTTGGCAGTGGAG	AATTTCAGTGGACTCCAGGAGGTCCGCTCG	708
181	ThrAspValAspAlaThrTrpGlnTrpLys	AsnPheSerGlyLeuGlnGluValArgSer	200

FIG. 4 (cont. next page)

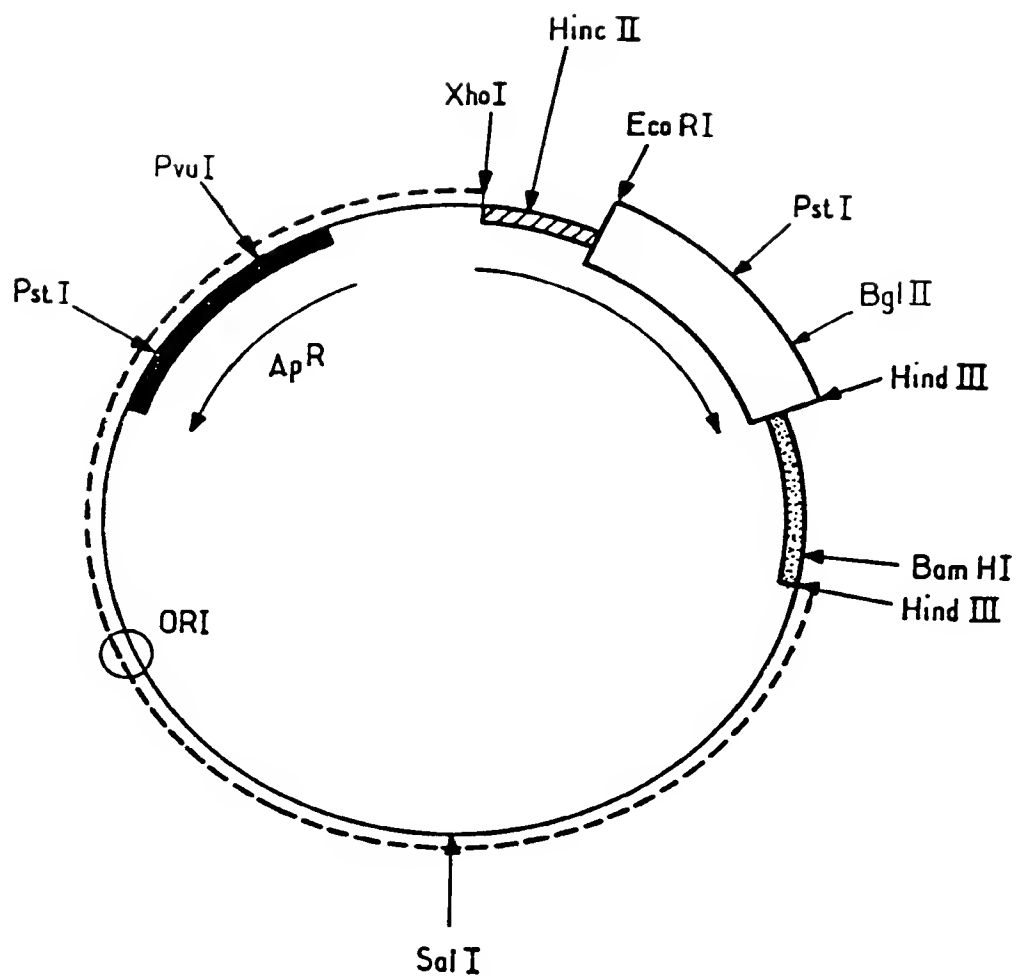
709	CACGTGCCCTAAGTTCGATGCTACCTGGGCC	ACTGCTCGCGAGGTCACTCTGAAGACTTTT	768
201	HisValProLysPheAspAlaThrTrpAla	ThrAlaArgGluValThrLeuLysThrPhe	220
	<u>T23</u>		
769	GCTGAAGATAACAGTGCAGCGTGCAGGCC	ACTATGTACAAGATGGCAGAGCAATCCTG	828
221	AlaGluAspAsnSerAlaSerValGlnAla	ThrMetTyrLysMetAlaGluGlnIleLeu	240
	<u>V2</u>		
829	GCGCGCCAGCAGCTGATCGAGACTGTCGAG	TACTCGTTGCCCTAACAAAGCACTATTTCGAA	888
241	AlaArgGlnGlnLeuIleGluThrValGlu	TyrSerLeuProAsnLysHisTyrPheGlu	260
	<u>V1</u>		
889	ATCGACCTGAGCTGGCACAAGGCCCTCCAA	AACACCGGCAAGACGCCGAGGTCTTCGCT	948
261	IleAspLeuSerTrpHisLysGlyLeuGln	AsnThrGlyLysAsnAlaGluValPheAla	280
	<u>T27</u>		
949	CCTCAGTCGGACCCCAACGGTCTGATCAAG	TGTACCGTCGGCCGGTCTCTCTGAAGTCT	1008
281	ProGlnSerAspProAsnGlyLeuIleLys	CysThrValGlyArgSerSerLeuLysSer	300
1009	AAATTGTAA		
301	LysLeuEnd		

## FIG. 4 (contd.)

DNA sequence opened by ATG in position 109 in Figure 3 and polypeptide coded for.

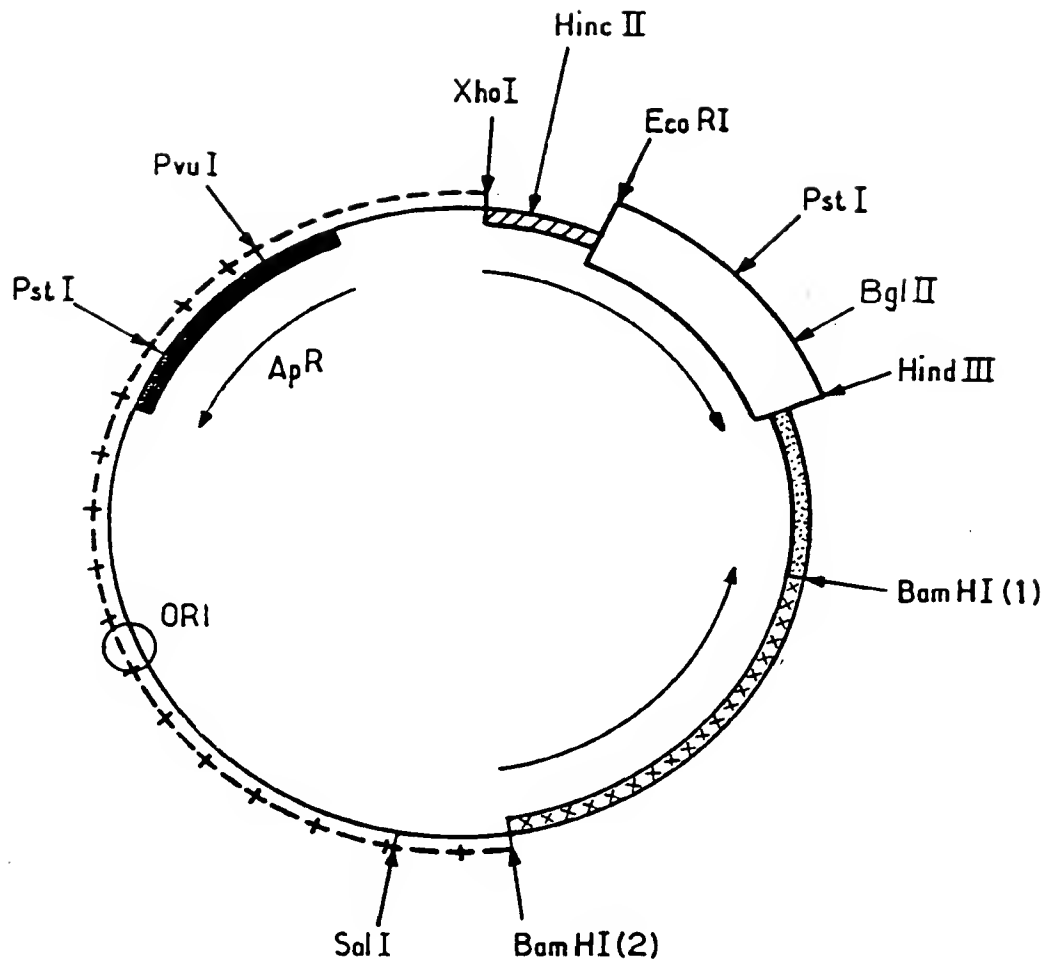
The sequenced peptides obtained by hydrolysis of *A. flavus* urate oxidase with trypsin and protease V8 are shown by arrows opposite the polypeptide coded for, according to

← : tryptic peptide  
 ← - - - : peptide obtained by hydrolysis with protease V8.



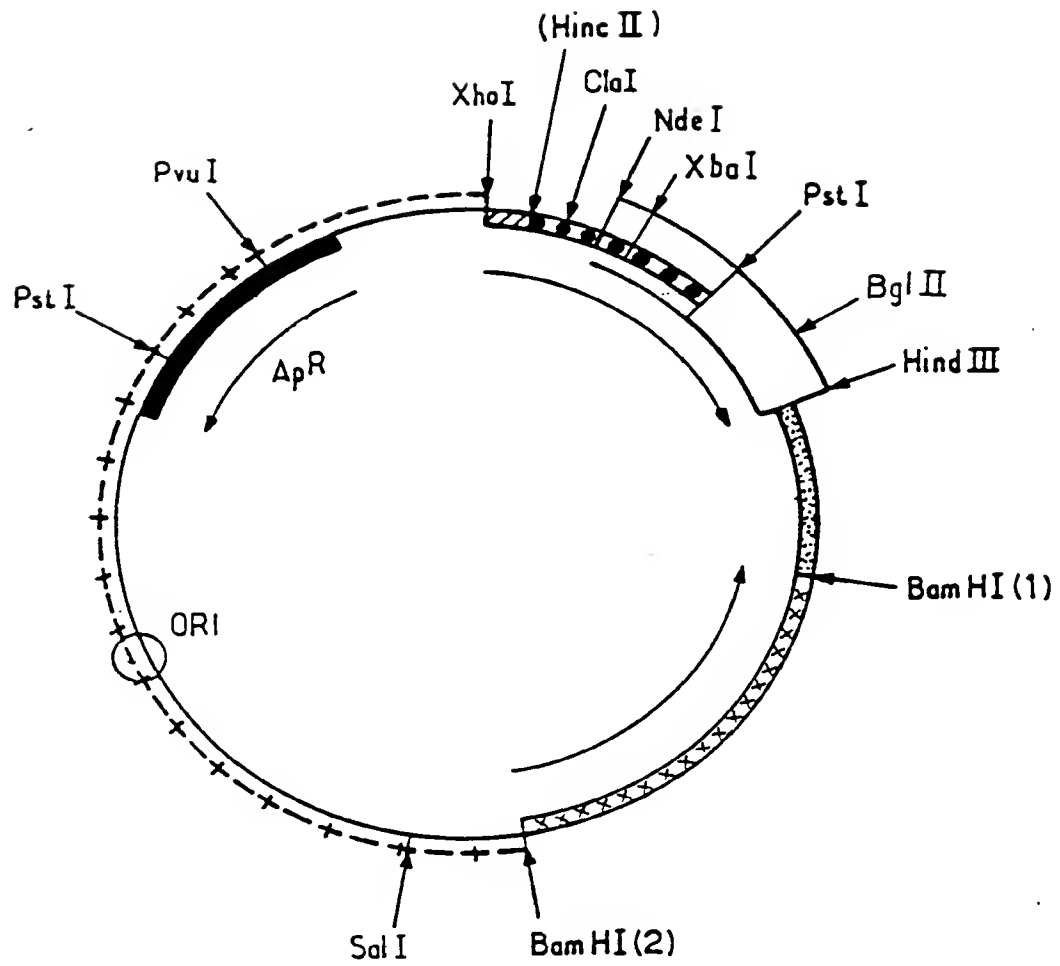
**FIG. 5**

*Plasmid p 163,1*



**FIG. 6**

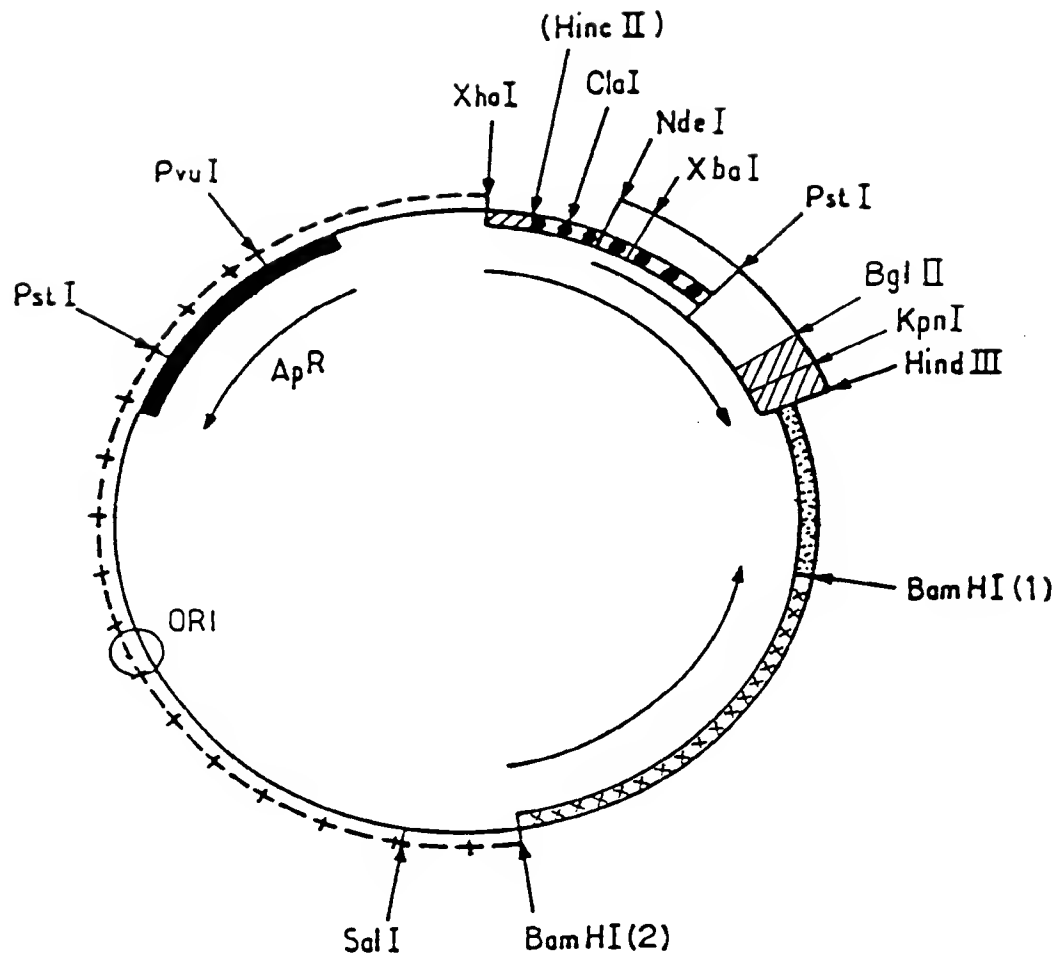
Plasmid p 160



**FIG. 7**

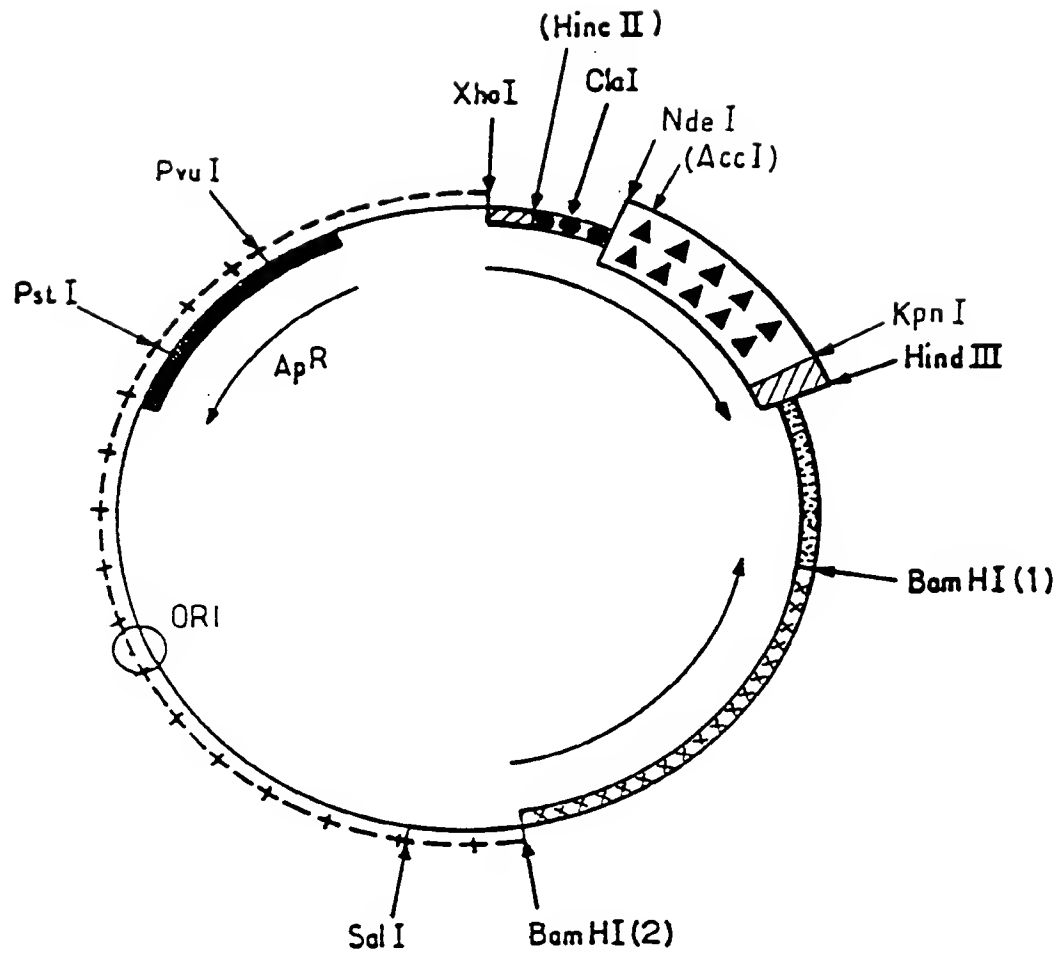
Plasmid p 373,2





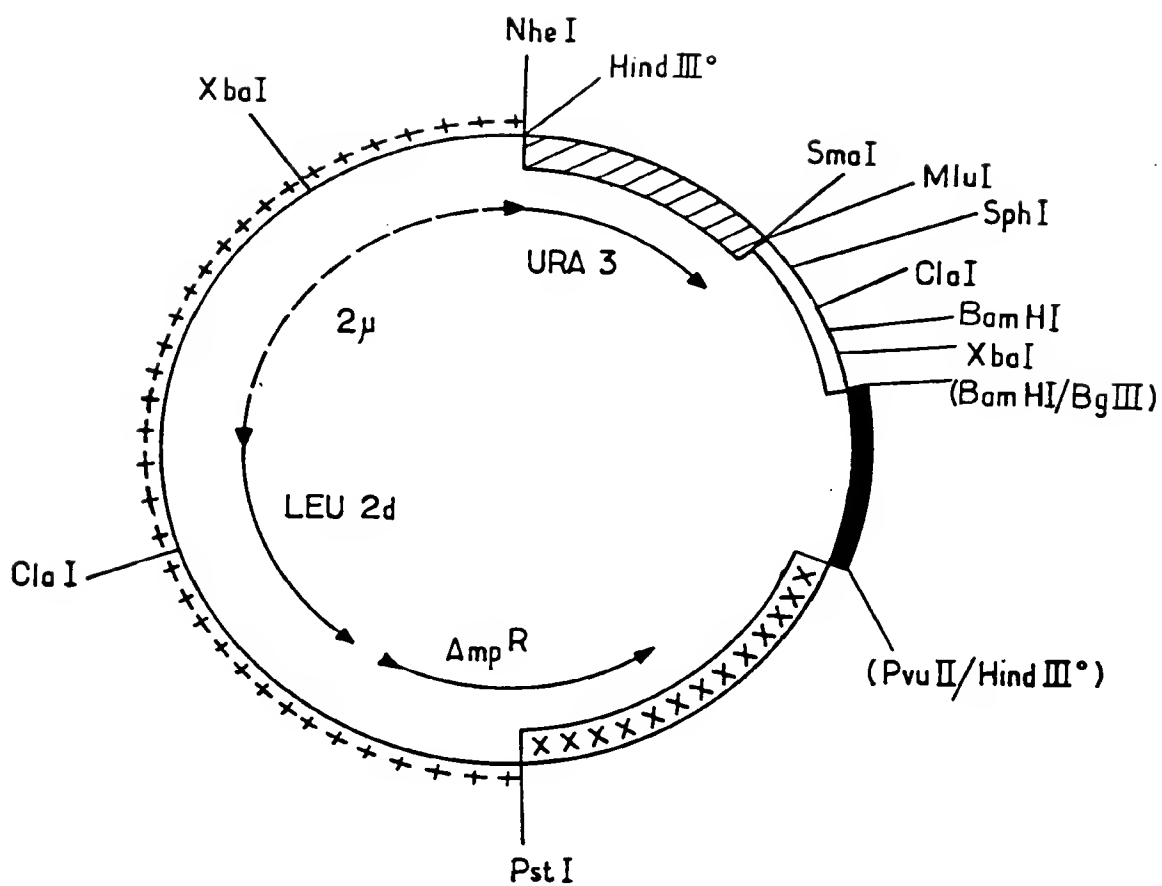
**FIG. 8**

Plasmid p 462



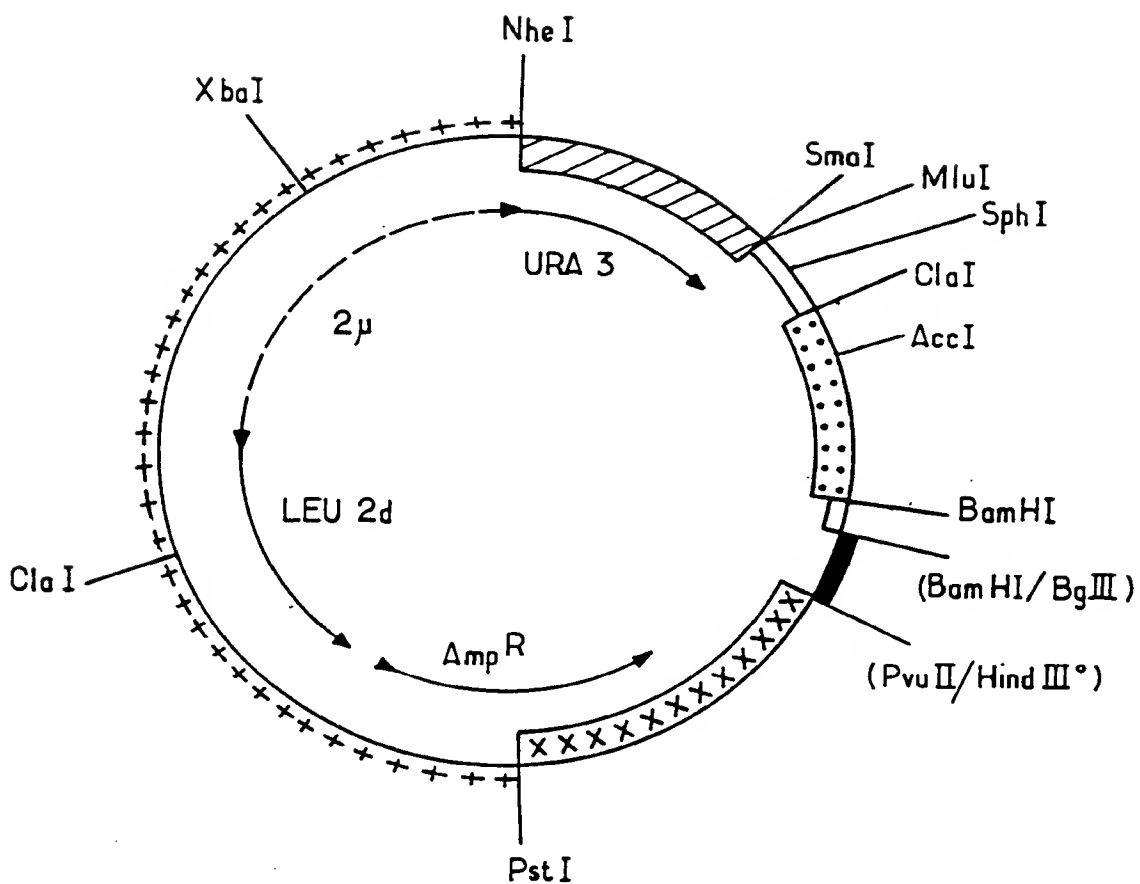
**FIG. 9**

Plasmid p466



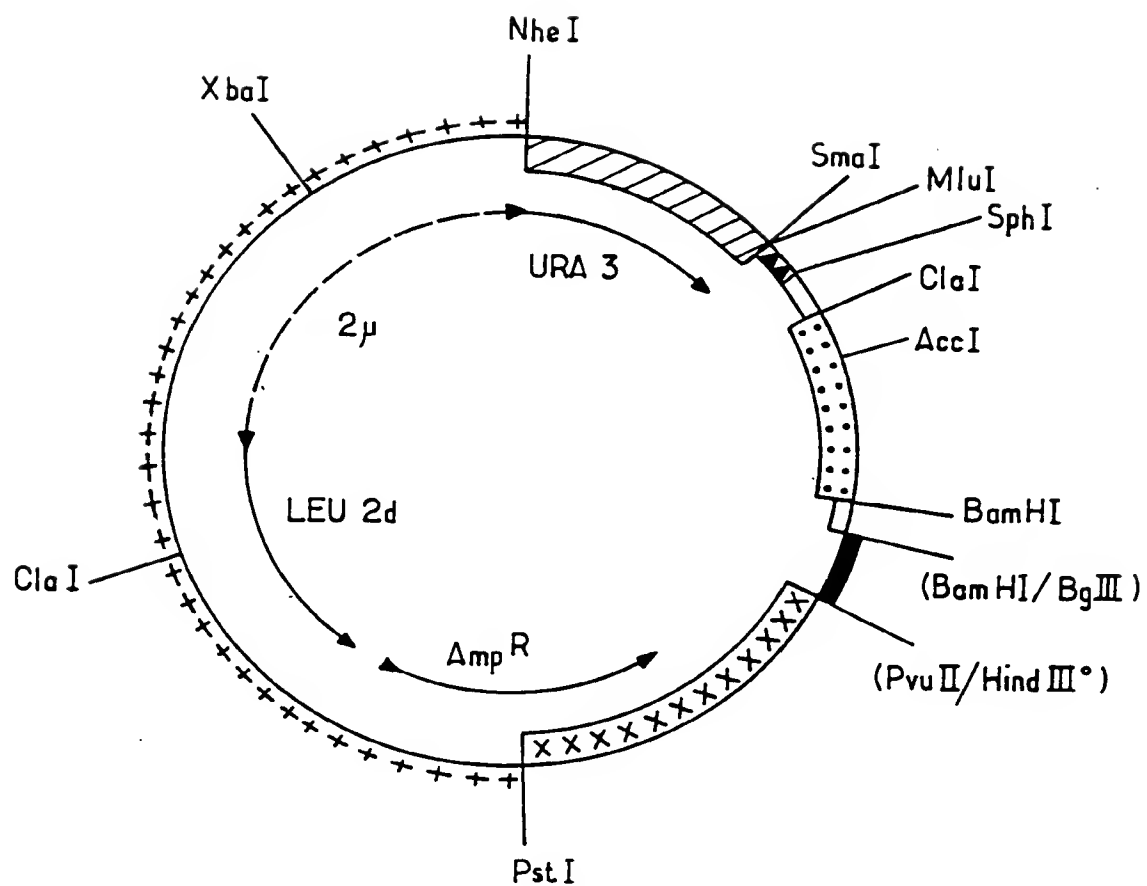
**FIG.10**

Plasmid pEMR 414



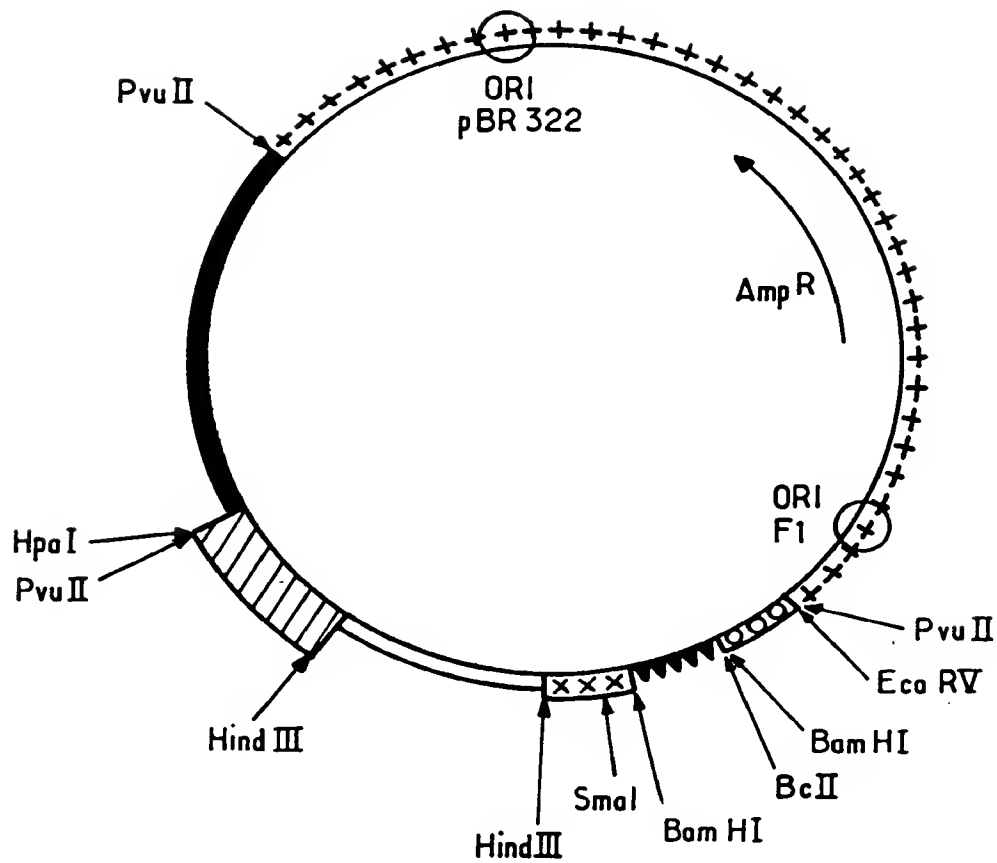
**FIG.11**

Plasmid pEMR 469



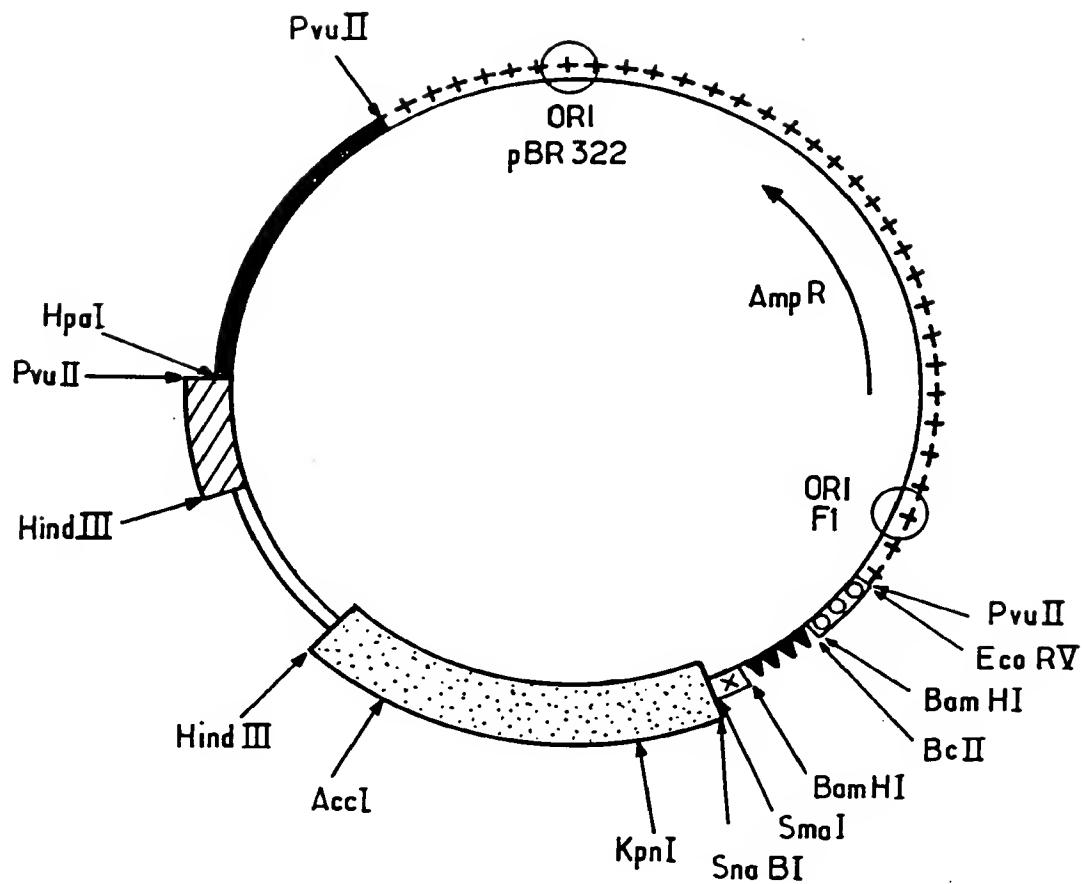
**FIG.12**

Plasmid pEMR 473



**FIG.13**

Plasmid PSE1



**FIG.14**

Plasmid pSV860